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Sequence Listing could not be accepted.

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217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: Mon May 21 14:19:56 EDT 2007

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Reviewer Comments:

<210> 1

<211> 660

<212> DNA

<213> Homo Sapiens Keratin 5

<400> 1

Invalid Response for <213>, It should have only Artificial, Unknown or
Genus species. This type of error is all through the sequences.

Application No: 10712629 Version No: 6.0

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Finished: 2007-05-17 19:43:02.403
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No. of SeqIDs Defined: 20
Actual SeqID Count: 20

SEQUENCE LISTING

<110> The Procter & Gamble Company

<120> Composition for Comprising a Mouse HRT Protein-Human Interacting
Partner Protein Complex (Revised)

<130> 9423

<140> 10712629

<141> 2003-11-13

<150> 10/712,629

<151> 2003-11-13

<160> 20

<170> PatentIn version 3.3

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<210> 2

<211> 746

<212> DNA

<213> Homo sapiens Ubiquitous Receptor

<400> 2

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<212> DNA

<213> Homo Sapiens Protein Inhibitor of Activated STAT-1

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<211> 792

<212> DNA

<213> Homo Sapiens Similar to Stromal Antigen 2

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<211> 747

<212> DNA

<213> Homo Sapiens Nucleoporin 160 Kda

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 <213> Homo Sapiens Retinoic Acid Receptor Gamma-1

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 <212> DNA
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 <213> Homo sapiens Annexin A1

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 <211> 323
 <212> DNA
 <213> Homo sapiens HIC Protein Isoform P32 and Isoform 40

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<210> 10

<211> 610

<212> DNA

<213> Homo Sapiens Insulin-like Growth Factor Binding Domain Protein 6

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<210> 11

<211> 718

<212> DNA

<213> Homo sapiens Inner Membrane Protein, Mitochondrial

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<212> DNA
<213> Homo Sapiens Endoplasmic reticulum thioredoxin superfamily member

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<211> 779
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<213> Homo Sapiens Protein Inhibitor of Activated STAT-3

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<211> 738

<212> DNA

<213> Homo Sapiens DEAD box polypeptide 3

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<210> 15

<211> 450

<212> DNA

<213> Homo Sapiens Dpy-30 Like Protein

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<210> 16
<211> 1269
<212> DNA
<213> Mus Musculus Vitamin D Receptor

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<210> 17

<211> 2079

<212> PRT

<213> Nucleotide sequence of HRT corresponding to the amino acid residue of the C-terminal portion of HR protein

<400> 17

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 1 5 10 15

Gly Cys Thr Gly Thr Gly Thr Cys Cys Ala Gly Gly Cys Ala Gly Cys
 20 25 30

Thr Gly Gly Ala Gly Ala Gly Gly Thr Ala Gly Gly Gly Gly Thr Ala
 35 40 45

Cys Thr Gly Ala Cys Cys Gly Gly Cys Cys Ala Cys Thr Cys Cys Cys
 50 55 60

Ala Gly Ala Ala Ala Thr Cys Ala Cys Gly Thr Ala Gly Gly Thr Cys
 65 70 75 80